RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/554,386_
Source: _PY/10_
Date Processed by STIC: _11/1/05_

ENTERED

PCT

RAW SEQUENCE LISTING   DATE: 11/01/2005
PATENT APPLICATION: US/10/554,386   TIME: 10:58:39

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      5 <120> TITLE OF INVENTION: GENES UPREGULATED IN A TOMATO PLANT HAVING AN INCREASED
      6       ANTHOCYANIN CONTENT PHENOTYPE
      8 <130> FILE REFERENCE: EP04-001C-US
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/554,386
C--> 10 <141> CURRENT FILING DATE: 2005-10-25
     10 <150> PRIOR APPLICATION NUMBER: US 60/465,605
     11 <151> PRIOR FILING DATE: 2003-04-25
     13 <160> NUMBER OF SEQ ID NOS: 12
     15 <170> SOFTWARE: PatentIn version 3.2
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     18 <211> LENGTH: 1675
     19 <212> TYPE: DNA
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     27 gaaaagtaca aaggattggt gggcaatttt ctgtgtggag acattgaaac tgtggagaat   180
     29 tggaggtcct atagcattca acataatctg ccaatatgga gtgaattctc ttacaaatat   240
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     33 tagtactttt tcttttggct tcatgatggg aatgggtagt gcattggaga cactatgtgg   360
     35 ccaggcatat ggagcagggc aagttcacat gttgggcgtt tatatgcaac gatccatcat   420
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     55 caatggatgg gagagcatgc ttttcattgg aattaatgct gccataagca tccgagtctc  1020
     57 aaatgaactg gggcaaggac atccaagagc taccaaatac tcggtctaca tcacggtgtt  1080
     59 tcagtccctc ctaatagggo tactctgcat ggtaattgta ctagtagcta gagatcatct  1140
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95 Asp Trp Trp Ala Ile Phe Cys Val Glu Thr Leu Lys Leu Trp Arg Ile
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99 Gly Gly Pro Ile Ala Phe Asn Ile Ile Cys Gln Tyr Gly Val Asn Ser
100     50                  55                  60
103 Leu Thr Asn Ile Phe Val Gly His Leu Gly Asn Val Glu Leu Ser Ala
104 65                  70                  75                  80
107 Ile Ser Ile Ala Gln Thr Val Ile Ser Thr Phe Ser Phe Gly Phe Met
108                 85                  90                  95
111 Met Gly Met Gly Ser Ala Leu Glu Thr Leu Cys Gly Gln Ala Tyr Gly
112                 100                 105                 110
115 Ala Gly Gln Val His Met Leu Gly Val Tyr Met Gln Arg Ser Ile Ile
116         115                 120                 125
119 Ile Leu Leu Ala Thr Cys Val Phe Leu Leu Pro Ile Tyr Leu Phe Thr
120     130                 135                 140
123 Thr Pro Leu Leu Val Leu Leu Gly Gln Glu Thr Ala Ile Ala Asp Leu
124 145                 150                 155                 160
127 Ser Gly Arg Tyr Thr Met Leu Leu Ile Pro Gln Leu Phe Ser Leu Ala
128                 165                 170                 175
131 Ile Asn Phe Pro Thr Ser Lys Phe Leu Gln Ala Gln Ser Lys Val Asp
132                 180                 185                 190
135 Val Leu Ala Gly Ile Gly Phe Ala Ala Val Leu Val His Ala Leu Phe
136         195                 200                 205
139 Leu Trp Leu Phe Ile Tyr Thr Leu Glu Trp Gly Thr Asn Gly Ala Ala
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147 Tyr Val Val Gly Trp Cys Lys Asp Gly Trp Lys Gly Leu Ser Trp Ser
148                 245                 250                 255
151 Ala Phe Asn Glu Ile Trp Ala Phe Val Arg Leu Ser Ile Ala Ser Ala
152                 260                 265                 270
155 Val Met Leu Cys Leu Glu Ile Trp Tyr Met Met Ser Ile Ile Leu Leu
156         275                 280                 285
159 Val Gly His Leu Asn Asn Ala Val Ile Ala Val Gly Ser Ile Ser Ile
160     290                 295                 300
163 Cys Met Asn Ile Asn Gly Trp Glu Ser Met Leu Phe Ile Gly Ile Asn
164 305                 310                 315                 320
167 Ala Ala Ile Ser Ile Arg Val Ser Asn Glu Leu Gly Gln Gly His Pro
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171 Arg Ala Thr Lys Tyr Ser Val Tyr Ile Thr Val Phe Gln Ser Leu Leu
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183 Leu Ala Tyr Leu Leu Gly Ile Thr Met Val Leu Asn Ser Val Gln Pro
184 385             390             395             400
187 Val Ile Ser Gly Val Ala Val Gly Gly Gly Trp Gln Ala Leu Val Ala
188             405             410             415
191 Tyr Ile Asn Leu Gly Cys Tyr Tyr Val Phe Gly Leu Pro Leu Gly Tyr
192         420             425             430
195 Leu Leu Gly Tyr Val Ala Lys Leu Gly Thr Lys Gly Leu Trp Leu Gly
196         435             440             445
199 Met Ile Ala Gly Ala Ala Leu Gln Thr Leu Leu Leu Leu Ile Ile Leu
200     450             455             460
203 Tyr Lys Thr Asn Trp Asn Lys Glu Val Asn Asp Thr Thr Glu Arg Met
204 465             470             475             480
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217 <212> TYPE: DNA
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225 accatggaag atgtcactgc caagattgat gcagttgaaa tcgactccaa gagtgttgtt    180
227 gctctgaagg agaaggcgat cgaaggtgaa gtaaagacta cagaaacaac acaagagaag    240
229 actaatgaag ctaaaagtaa tccggagaaa gaatccaatg gtgcaacttt caaagaagag    300
231 gaagttccag ttgaggttga acctaaaact ggagtctctt ttcctgtcaa acttgaagat    360
233 ggaaagcagt tgaaagctgt aggtttaagg aagaagagca tgcttggcat gggcatcaaa    420
235 atctatggct ttggaatata tgcagacaat gagaaattga aagacctcat gcagtccaag    480
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251 gtagtaggtt ccagcatttt actagtctag gttgttaaag aataagctgt ggtgactttg    960
253 aactgatgaa taatagctca ataaaagcat ggttgatttt caaaaaaaaa aaaaaaaaa    1020
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278      50                  55                  60
281 Val Pro Val Glu Val Glu Pro Lys Thr Gly Val Ser Phe Pro Val Lys
282 65                  70                  75                  80
285 Leu Glu Asp Gly Lys Gln Leu Lys Ala Val Gly Leu Arg Lys Lys Ser
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341 tctgtcgaca aagttgaagc cagcattgac gcttctctcc ttaaatgcag aaaattaccc   240
343 tcaggatgca ttctacaaga acaatccaat gcacattgca aggtgacgtg ggtggaacac   300
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/554,386

DATE: 11/01/2005
TIME: 10:58:39

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Output Set: N:\CRF4\11012005\J554386.raw

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Please Note:
**Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.**

Seq#:11; N Pos. 712,713,714,752,758,801,811,827

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L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
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